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## RAW SEQUENCE LISTING

DATE: 02/26/2003

PATENT APPLICATION: US/09/077,173D

TIME: 14:49:36

Input Set : A:\Sequence.txt

Output Set: N:\CRF4\02262003\I077173D.raw

3 <110> APPLICANT: EUROSCREEN S.A.  
4 COMMUNI, DIDIER  
5 PIROTON , SABINE  
6 PARMENTIER, MARC  
7 BOEYNAEMS, JEAN-MARIE  
9 <120> TITLE OF INVENTION: HUMAN PYRIMIDINE RECEPTOR  
11 <130> FILE REFERENCE: 9409/2112  
13 <140> CURRENT APPLICATION NUMBER: US 09/077,173D  
14 <141> CURRENT FILING DATE: 1998-11-12  
16 <150> PRIOR APPLICATION NUMBER: PCT/BE96/00123  
17 <151> PRIOR FILING DATE: 1996-11-21  
19 <150> PRIOR APPLICATION NUMBER: EPO 95870124.5  
20 <151> PRIOR FILING DATE: 1995-11-21  
22 <160> NUMBER OF SEQ ID NOS: 4  
24 <170> SOFTWARE: PatentIn version 3.2  
26 <210> SEQ ID NO: 1  
27 <211> LENGTH: 1429  
28 <212> TYPE: DNA  
29 <213> ORGANISM: Homo sapiens  
31 <400> SEQUENCE: 1

ENTERED

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36 gggggtgggc agggaaatcc tgccaccctc acttctcccc ttcccatctc cagggggggcc      180
38 atggccagta cagagtcctc cctgttgaga tccctaggcc tcagcccagg tcctggcagc      240
40 agtgaggtag agctggactg ttggtttgat gaggatttca agttcatcct gctgcctgtg      300
42 agctatgcag ttgtctttgt gctgggcttg ggccttaacg cccaaccct atggctcttc      360
44 atcttccgcc tccgaccctg ggatgcaacg gccacctaca tgttccacct ggcattgtca      420
46 gacaccttgt atgtgtgtgc gctgcccacc ctcatctact attatgcagc ccacaaccac      480
48 tggccctttg gcaactgagat ctgcaagttc gtccgctttc ttttctattg gaacctctac      540
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52 cttcgggcac tacgtgagg ccgcccctgc ctgcaggcc ttctctgctt ggcaagtttg      660
54 ttggtcgtag ccggtgcct cgtgcccac ctgttctttg tcacaaccag caacaaagg      720
56 accaccgtcc tgtgccatga caccactcgg cctgaagagt ttgaccacta tgtgcacttc      780
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60 ggactcatgg ctgcgcct gtatcagccc ttgccaggct ctgcacagtc gtcttctcgc      900
62 ctccgctctc tccgcaccat agctgtggtg ctgactgtct ttgctgtctg cttcgtgcct      960
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66 ctgaacattg tcaacgtggt ctataaagt actcggcccc tggccagtgc caacagctgc      1080
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70 tgtggtggtg gcaagcccca gcccgcacg gctgcctctt ccctggcact agtgtccctg      1200
72 cctgaggata gcagctgcag gtgggcggcc acccccagg acagtagctg ctctactcct      1260
74 agggcagata gattgtaaca cgggaagccg gcaagtgaga gaaaagggga tgagtgcagg      1320
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81 <210> SEQ ID NO: 2
82 <211> LENGTH: 365
83 <212> TYPE: PRT
84 <213> ORGANISM: Homo sapiens
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88 Met Ala Ser Thr Glu Ser Ser Leu Leu Arg Ser Leu Gly Leu Ser Pro
89 1          5          10          15
92 Gly Pro Gly Ser Ser Glu Val Glu Leu Asp Cys Trp Phe Asp Glu Asp
93          20          25          30
96 Phe Lys Phe Ile Leu Leu Pro Val Ser Tyr Ala Val Val Phe Val Leu
97          35          40          45
100 Gly Leu Gly Leu Asn Ala Pro Thr Leu Trp Leu Phe Ile Phe Arg Leu
101          50          55          60
104 Arg Pro Trp Asp Ala Thr Ala Thr Tyr Met Phe His Leu Ala Leu Ser
105 65          70          75          80
108 Asp Thr Leu Tyr Val Leu Ser Leu Pro Thr Leu Ile Tyr Tyr Tyr Ala
109          85          90          95
112 Ala His Asn His Trp Pro Phe Gly Thr Glu Ile Cys Lys Phe Val Arg
113          100         105         110
116 Phe Leu Phe Tyr Trp Asn Leu Tyr Cys Ser Val Leu Phe Leu Thr Cys
117          115         120         125
120 Ile Ser Val His Arg Tyr Leu Gly Ile Cys His Pro Leu Arg Ala Leu
121          130         135         140
124 Arg Trp Gly Arg Pro Arg Leu Ala Gly Leu Leu Cys Leu Ala Val Trp
125 145         150         155         160
128 Leu Val Val Ala Gly Cys Leu Val Pro Asn Leu Phe Phe Val Thr Thr
129          165         170         175
132 Ser Asn Lys Gly Thr Thr Val Leu Cys His Asp Thr Thr Arg Pro Glu
133          180         185         190
136 Glu Phe Asp His Tyr Val His Phe Ser Ser Ala Val Met Gly Leu Leu
137          195         200         205
140 Phe Gly Val Pro Cys Leu Val Thr Leu Val Cys Tyr Gly Leu Met Ala
141          210         215         220
144 Arg Arg Leu Tyr Gln Pro Leu Pro Gly Ser Ala Gln Ser Ser Ser Arg
145 225          230         235         240
148 Leu Arg Ser Leu Arg Thr Ile Ala Val Val Leu Thr Val Phe Ala Val
149          245         250         255
152 Cys Phe Val Pro Phe His Ile Thr Arg Thr Ile Tyr Tyr Leu Ala Arg
153          260         265         270
156 Leu Leu Glu Ala Asp Cys Arg Val Leu Asn Ile Val Asn Val Val Tyr
157          275         280         285
160 Lys Val Thr Arg Pro Leu Ala Ser Ala Asn Ser Cys Leu Asp Pro Val
161          290         295         300
164 Leu Tyr Leu Leu Thr Gly Asp Lys Tyr Arg Arg Gln Leu Arg Gln Leu
165 305         310         315         320
168 Cys Gly Gly Gly Lys Pro Gln Pro Arg Thr Ala Ala Ser Ser Leu Ala
169          325         330         335
172 Leu Val Ser Leu Pro Glu Asp Ser Ser Cys Arg Trp Ala Ala Thr Pro

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176 Gln Asp Ser Ser Cys Ser Thr Pro Arg Ala Asp Arg Leu
177          355          360          365
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181 <211> LENGTH: 35
182 <212> TYPE: DNA
183 <213> ORGANISM: Homo sapiens
185 <400> SEQUENCE: 3
186 cagatctaga tactatgttc tacactctta cgtgc 35
189 <210> SEQ ID NO: 4
190 <211> LENGTH: 35
191 <212> TYPE: DNA
192 <213> ORGANISM: Homo sapiens
194 <400> SEQUENCE: 4
195 tcttaagctt ggagtcacgt acgagcaagc tagtt 35
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**VERIFICATION SUMMARY**

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